

Huynh

#20



1600

RAW SEQUENCE LISTING

DATE: 06/26/2003

PATENT APPLICATION: US/09/712,819C

TIME: 11:04:19

Input Set : A:\US09712819C.raw

Output Set: N:\CRF4\06262003\I712819C.raw

SEQUENCE LISTING

1 (1) GENERAL INFORMATION:
2 (i) APPLICANT: Stevens, Fred J.
3 Argon, Yair
4 Davis, David P.
5 Raffen, Rosemarie
6 (ii) TITLE OF INVENTION: Fibril-Blocking Peptide, A Method for Preventing
7 Fibril Formation
8 (iii) NUMBER OF SEQUENCES: 11
9 (iv) CORRESPONDENCE ADDRESS:
10 (A) ADDRESSEE: CHERSKOV & FLAYNIK
11 (B) STREET: 20 N. Wacker Drive
12 (C) CITY: Chicago
13 (D) STATE: Illinois
14 (E) COUNTRY: United States
15 (F) ZIP: 60606
16 (v) COMPUTER READABLE FORM:
17 (A) MEDIUM TYPE: 3.50 inch, 1.4 MB storage
18 (B) COMPUTER: PC
19 (C) OPERATING SYSTEM: Microsoft Windows 98
20 (D) SOFTWARE: Wordperfect
21 (vi) CURRENT APPLICATION DATA:
C--> 22 (A) APPLICATION NUMBER: US/09/712,819C
C--> 23 (B) FILING DATE: 11-Mar-2002
24 (vii) PRIOR APPLICATION DATA:
25 (A) APPLICATION NUMBER: U.S. 60/165,424
26 (B) FILING DATE: 1999-NOV-14
27 (viii) ATTORNEY/AGENT INFORMATION:
28 (A) NAME: Cherskov, Michael J.
29 (B) REGISTRATION NUMBER: 33,664
30 (C) REFERENCE/DOCKET NUMBER: 0003/00537
31 (ix) TELECOMMUNICATION INFORMATION:
32 (A) TELEPHONE: (312) 621-1330
33 (B) TELEFAX: (312) 621-0088
34 (2) INFORMATION FOR SEQ ID NO: 1:
35 (i) SEQUENCE CHARACTERISTICS:
36 (A) LENGTH: 7 amino acids
37 (B) TYPE: amino acid
38 (C) STRANDEDNESS: Single
39 (D) TOPOLOGY: linear
40 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
41 Phe Thr Leu Thr Ile Ser Ser
42 1 5

ENTERED

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44 (2) INFORMATION FOR SEQ ID NO: 2:
45 (i) SEQUENCE CHARACTERISTICS:
46 (A) LENGTH: 7 amino acids
47 (B) TYPE: amino acid
48 (C) STRANDEDNESS: Single
49 (D) TOPOLOGY: linear
50 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
51 Thr Arg Ile Phe Ser Lys Leu
52 1 5

54 (2) INFORMATION FOR SEQ ID NO: 3:
55 (i) SEQUENCE CHARACTERISTICS:
56 (A) LENGTH: 7 amino acids
57 (B) TYPE: amino acid
58 (C) STRANDEDNESS: Single
59 (D) TOPOLOGY: linear
60 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
61 Phe Tyr Gln Leu Ala Leu Thr
62 1 5

64 (2) INFORMATION FOR SEQ ID NO: 4:
65 (i) SEQUENCE CHARACTERISTICS:
66 (A) LENGTH: 8 amino acids
67 (B) TYPE: amino acid
68 (C) STRANDEDNESS: Single
69 (D) TOPOLOGY: linear
70 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
71 Pro Lys Leu Leu Ile Tyr Trp Ala
72 1 5

74 (2) INFORMATION FOR SEQ ID NO: 5:
75 (A) LENGTH: 7 amino acids
76 (B) TYPE: amino acid
77 (C) STRANDEDNESS: Single
78 (D) TOPOLOGY: linear
79 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
80 Thr Asp Phe Thr Leu Thr Ile
81 1 5

83 (2) INFORMATION FOR SEQ ID NO: 6:
84 (i) SEQUENCE CHARACTERISTICS:
85 (A) LENGTH: 7 amino acids
86 (B) TYPE: amino acid
87 (C) STRANDEDNESS: Single
88 (D) TOPOLOGY: linear
89 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
90 Phe Thr Leu Lys Ile Ser Arg
91 1 5

93 (2) INFORMATION FOR SEQ ID NO: 7:
94 (i) SEQUENCE CHARACTERISTICS:
95 (A) LENGTH: 9 amino acids
96 (B) TYPE: amino acid
97 (C) STRANDEDNESS: Single

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98         (D) TOPOLOGY: linear
99         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
100 Thr Asp Phe Thr Leu Thr Ile Ser Ser
101 1             5
103 (2) INFORMATION FOR SEQ ID NO: 8
104     (i) SEQUENCE CHARACTERISTICS:
105         (A) LENGTH: 11 amino acids
106         (B) TYPE: amino acid
107         (C) STRANDEDNESS: Single
108         (D) TOPOLOGY: linear
109     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
110 Tyr Gly Arg Lys Lys Arg Arg Gln Lys Lys Lys
111 1             5             10
113 (2) INFORMATION FOR SEQ ID NO: 9
114     (i) SEQUENCE CHARACTERISTICS:
115         (A) LENGTH: 29 amino acids
116         (B) TYPE: amino acid
117         (C) STRANDEDNESS: Single
118         (D) TOPOLOGY: linear
119     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
120 Tyr Glu Arg Lys Lys Arg Arg Gln Arg Arg Arg Ser Gly Ser Gly
121 1             5             10             15
122 Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala Glu Asp
123             20             25
125 (2) INFORMATION FOR SEQ ID NO: 10
126     (i) SEQUENCE CHARACTERISTICS:
127         (A) LENGTH: 29 amino acids
128         (B) TYPE: amino acid
129         (C) STRANDEDNESS: Single
130         (D) TOPOLOGY: linear
131     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
132 Tyr Glu Arg Lys Lys Arg Arg Gln Arg Arg Arg Ser Gly Ser Gly
133 1             5             10             15
134 Thr Asp Ala Thr Ala Pro Ala Ser Ser Leu Gln Ala Glu Asp
135             20             25
137 (2) INFORMATION FOR SEQ ID NO: 11:
138     (i) SEQUENCE CHARACTERISTICS:
139         (A) LENGTH: 4 amino acids
140         (B) TYPE: amino acid
141         (C) STRANDEDNESS: Single
142         (D) TOPOLOGY: linear
143     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
144 Ala Leu Leu Asn
145 1

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VERIFICATION SUMMARY

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L:22 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:23 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]